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* OM nucleic - nucleic search, using sw model

Run on: January 24, 2002, 21:47:46 ; Search time 130.07 Seconds
 Sequence: (without alignments) 2287.938 Million cell updates/sec

Title: US-09-464-528-6

Perfect score: 1314

Sequence: ttcttagatcaaactacatcc.....tttttgaagataaccatgg 1314

Scoring table: IDENTITY.NUC Gapop 10.0 , Gapext 1.0

Searched: 351203 seqs, 113238999 residues

Total number of hits satisfying chosen parameters: 702406

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Issued_Parents.NA:*

1: /cgn2_6/ptodata/2/ina/5A_COMB_seq:*

2: /cgn2_6/ptodata/2/ina/5B_COMB_seq:*

3: /cgn2_6/ptodata/2/ina/6A_COMB_seq:*

4: /cgn2_6/ptodata/2/ina/6B_COMB_seq:*

5: /cgn2_6/ptodata/2/ina/PCITS_COMB_seq:*

6: /cgn2_6/ptodata/2/ina/backfiles1.seq:*

Score: No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match Length	DB ID	Description	
1	78	5.9	7218	1 US-08-232-463-14	
2	61.4	837	4 US-08-232-463-14	Sequence 14, APPL	
3	56.2	4.3	4 US-08-232-463-14	Sequence 14, APPL	
4	56	4.3	19124	2 US-08-407-8269-13	Sequence 13, APPL
5	55.4	615	4 US-08-998-416-1137	Sequence 13, APPL	
6	54.4	4.1	837	4 US-08-998-416-1137	Sequence 13, APPL
7	54.4	4.1	186	2 US-08-998-416-1137	Sequence 13, APPL
8	54.2	4.1	186	2 US-08-998-416-1137	Sequence 13, APPL
9	53.6	4.1	19124	2 US-08-487-8228-13	Sequence 13, APPL
10	52.6	4.0	665	2 US-08-883-7958-36	Sequence 13, APPL
11	52.2	4.0	1826	4 US-09-286-691-11	Sequence 11, APPL
12	52.2	4.0	1826	4 US-09-687-147-11	Sequence 11, APPL
13	52	4.0	8920	2 US-08-440-8550-1	Sequence 1, APPL
14	52	4.0	8920	4 US-09-150-741-1	Sequence 1, APPL
15	51.2	3.9	3933	2 US-08-731-722-3	Sequence 3, APPL
16	51.2	3.9	3933	2 US-08-731-722-3	Sequence 3, APPL
17	51	3.9	615	4 US-08-998-416-1136	Sequence 186, APPL
18	51	3.9	665	2 US-08-883-7958-36	Sequence 36, APPL
19	50.8	602	1 US-08-222-393-1	Sequence 1, APPL	
20	50.2	3.8	473	1 US-08-764-100-14	Sequence 15, APPL
21	50.2	3.8	4970	1 US-08-764-100-14	Sequence 14, APPL
22	50.2	3.8	4970	1 US-08-764-100-20	Sequence 20, APPL
23	50.2	3.8	8920	2 US-08-440-8550-1	Sequence 1, APPL
24	49.8	4457	1 US-08-150-741-1	Sequence 1, APPL	
25	49.8	3.8	4457	2 US-08-910-553-1	Sequence 1, APPL
26	49.8	4467	2 US-08-909-425A-1	Sequence 1, APPL	
27	49.8	4467	2 US-08-909-425A-1	Sequence 1, APPL	

RESULT

RESULT	1
US-08-232-463-14	
Sequence 14, Application	US/08232463
Patent No.	5670367
GENERAL INFORMATION:	
APPLICANT:	DORNER, F.
APPLICANT:	SCHEIFLINGER, F.
APPLICANT:	FAIRNER, F. G.
TITLE OF INVENTION:	RECOMBINANT FOWLPOX VIRUS
NUMBER OF SEQUENCES:	52
CORRESPONDENCE ADDRESS:	
ADDRESSEE:	Foley & Lardner
STREET:	1800 Diagonal Road, Suite 500
CITY:	Alexandria
STATE:	VA
ZIP:	22313-0299
IP:	22313-0299
COMPUTER READABLE FORM:	
COMPUTER:	IBM PC compatible
OPERATING SYSTEM:	PC-DOS/MS-DOS
SOFTWARE:	Patentin Release #1.0, Version
CURRENT APPLICATION DATA:	
APPLICATION NUMBER:	US/08/232,463
FILING DATE:	
CLASSIFICATION:	435
PRIOR APPLICATION DATA:	
APPLICATION NUMBER:	US/07/935,313
FILING DATE:	
APPLICATION NUMBER:	EP 91 114 300.6
FILING DATE:	26-AUG-1991
ATTORNEY/AGENT INFORMATION:	
NAME:	BENT, Stephen A.
REGISTRATION NUMBER:	29,768
REFERENCE/DOCKET NUMBER:	30472/114 IMMU
TELECOMMUNICATION INFORMATION:	
TELEPHONE:	(703) 683-9300
TELEFAX:	(703) 683-4109
TELEFAX:	899149
INFORMATION FOR SEQ ID NO:	14:
SEQUENCE CHARACTERISTICS:	
LENGTH:	7218 base Pairs
TYPE:	nucleic acid
STRANDEDNESS:	single
TOPOLOGY:	linear
IMMEDIATE SOURCE:	
CLONE:	FIZ9pt-FIS
US-08-232-463-14	

Patent No. 5,213,972
Sequence 14, Appl
Sequence 32, Appl
Sequence 4, Appl
Sequence 1, Appl
Sequence 3, Appl
Sequence 1, Appl
Sequence 36, Appl
Sequence 35, Appl
Sequence 36, Appl
Sequence 36, Appl

TELEFAX: 919-541-8689
; INFORMATION FOR SEQ ID NO: 288:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 837 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
; ORIGINAL SOURCE:
; ORGANISM: PAG1241RP
US-08-998416-288

RESULT
US-08-998-416-288/C
Sequence 289 Application US/08998416
PATENT NO. 6239264
GENERAL INFORMATION:
APPLICANT: Philippson, Peter
APPLICANT: Poltmann, Rainer
APPLICANT: Steiner, Sabine
APPLICANT: Mohr, Christine
APPLICANT: Wenzel, Jürgen
APPLICANT: Knechtli, Philipp
APPLICANT: Rehischung, Corinne
TITLE OF INVENTION: GENOMIC DNA SEQUENCES OF ASHBYA GOSSYPII
TITLE OF INVENTION: AND USES THEREOF
NUMBER OF SEQUENCES: 1152
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6239264artis Corporation
STREET: 3054 Cornwallis Road
CITY: Research Triangle Park
STATE: No. 6239264th Carolina
COUNTRY: USA
ZIP: 27709
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/998,416
FILING DATE: 24-DEC-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: CH 0016/97
APPLICATION NUMBER: CH 0016/97
FILING DATE: 31-DEC-1996
ATTORNEY/AGENT INFORMATION:
NAME: Meigs, J. Timothy
REGISTRATION NUMBER: 38,241
REFERENCE/DOCKET NUMBER: PF/5-30306/A/GCG1976
TELECOMMUNICATION INFORMATION:
TELEPHONE: 919-541-8587

3
LT 3
8-998416-1137/C
quence 1137, Application US/08998416
ent No. 523964
GENERAL INFORMATION:
APPLICANT: Philippson, Peter
APPLICANT: Pohlmann, Rainer
APPLICANT: Steiner, Sabine
APPLICANT: Mohr, Christine
APPLICANT: Wendland, Jurgen
APPLICANT: Knechtel, Philipp
APPLICANT: Reblischung, Corinne
TITLE OF INVENTION: GENOMIC DNA SEQUENCES OF ASHYA GOSSYPII
TITLE OF INVENTION: AND USES THEREOF
NUMBER OF SEQUENCES: 1152
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 623964artis Corporation
STREET: 3054 Cornwallis Road
CITY: Research Triangle Park
STATE: NC 27709 USA
COUNTRY: USA
ZIP: 27709
COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/998,416
FILING DATE: 24-DEC-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: CH 0015/97
FILING DATE: 31-DEC-1996
ATTORNEY/AGENT INFORMATION:
NAME: Meigs, J. Timothy
REGISTRATION NUMBER: 38,241
REFERENCE/DOCKET NUMBER: PF15-30306/A/CGC1976
TELECOMMUNICATION INFORMATION:

TELEPHONE: 919-541-9587
TELEFAX: 919-541-8689
INFORMATION FOR SBO ID NO: 113
SEQUENCE CHARACTERISTICS:
LENGTH: 636 base Pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
ORIGINAL SOURCE:
ORGANISM: PAG1692RP
US-08-998416-1137

SEQUENCE CHARACTERISTICS:
 LENGTH: 19124 base Pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: cDNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
US-08-487-826B-13

US 08-487-826B-13
Sequence 13, Application US/08487826B
Patent No. 593627
GENERAL INFORMATION:
APPLICANT: Sim, Kim L.
APPLICANT: Chittim, Chetan
APPLICANT: Miller, Louis H.
APPLICANT: Peterson, David S.
APPLICANT: Su, Xin-zhuan
APPLICANT: Wellens, Thomas E.
TITLE OF INVENTION: BINDING DOMAIN
TITLE OF INVENTION: AND PLASMIDIUM
NUMBER OF SEQUENCES: 45
CORRESPONDENCE ADDRESS:
ADDRESSEE: Knobbe Martens Olson
SUBJECT: 620 Newport Center Drive
CITY: Newport Beach
STATE: California
COUNTRY: US
ZIP: 92660
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/487,82
FILING DATE: 10-SEP-1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Israelsen, Ned
REGISTRATION NUMBER: 29, 655
REFERENCE/DOCKET NUMBER: NIH121.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 235-8550
TELEFAX: (619) 235-0176
INFORMATION FOR SRO/TM NO: 13

RESULT 5 US-08-998-416-186
Sequence 186, Application US/08998416
Patent No. 6239264
GENERAL INFORMATION:
APPLICANT: Philipsen, Peter
APPLICANT: Pöhlmann, Rainer
APPLICANT: Steiner, Sabine
APPLICANT: Mohr, Christine
APPLICANT: Wendland, Jürgen
APPLICANT: Knechtli, Philipp
APPLICANT: Rebischung, Corinne
TITLE OF INVENTION: GENOMIC DNA SEQUENCES OF ASHYA G
TITLE OF INVENTION: AND USES THEREOF
NUMBER OF SEQUENCES: 1152
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6239264 artis Corporation
STREET: 3054 Cornwallis Road
CITY: Research Triangle Park
STATE: No. 6239264th Carolina
COUNTRY: USA
ZIP: 27709
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent Release #1.0, version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/998,416
FILING DATE: 24-DEC-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: CH 0016/97

Fri Jan 25 09:41:37 2002

us-09-464-528-6.rni

OM nucleic - nucleic search, using sw model

Run on: January 24, 2002, 22:04:33 ; Search time 130.07 Seconds
(without alignments)

Scoring table: IDENTITY.NUC

Gapop 10.0 , Gapext 1.0

Searched: 351203 seqs, 113238999 residues

Total number of hits satisfying chosen parameters: 702406

minimum DB seq length: 0

maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

database : Issued_Patents_NA.*

1: /cgn2_6/podata/2/ina/5A_Comb.seq:*

2: /cgn2_6/podata/2/ina/5B_Comb.seq:*

3: /cgn2_6/podata/2/ina/6A_Comb.seq:*

4: /cgn2_6/podata/2/ina/6B_Comb.seq:*

5: /cgn2_6/podata/2/ina/PCNUS_Comb.seq:*

6: /cgn2_6/podata/2/ina/backfiles1.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

RESULTS

Result No.	Score	Query	Match Length	DB	ID	Description
1	78	3.6	7218	1	US-08-232-463-14	Sequence 14, Appl
2	61.4	2.8	837	4	US-08-998-416-288	Sequence 288, App
3	57.2	3.0	3095	6	S231168-1	Patent No. 567367
4	57	2.6	837	4	US-08-998-416-288	Sequence 288, App
5	56.4	2.6	615	4	US-08-998-16-1137	Sequence 186, App
6	56.2	2.6	636	4	US-08-998-16-1137	Sequence 186, App
7	56	2.6	19124	2	US-08-4-826B-13	Sequence 13, Appl
8	54.8	2.5	636	4	US-08-998-16-1137	Sequence 1137, Appl
9	54.2	2.5	1186	2	US-08-7-311-722-5	Sequence 5, Appl
10	53.8	2.5	6243	2	US-09-056-075-1	Sequence 1, Appl
11	53.6	2.5	19124	2	US-08-887-826B-13	Sequence 13, Appl
12	53.2	2.5	3138	1	US-07-867-106-4	Sequence 4, Appl
13	52.6	2.4	665	2	US-08-883-795A-36	Sequence 36, Appl
14	52.2	2.4	1826	4	US-09-686-601-11	Sequence 11, Appl
15	52.2	2.4	1826	4	US-09-687-47-11	Sequence 11, Appl
16	52	2.4	8920	2	US-08-446-855A-1	Sequence 1, Appl
17	52	2.4	8920	4	US-09-150-741-1	Sequence 1, Appl
18	51.8	2.4	782	4	US-09-007-119-15	Sequence 15, Appl
19	51.2	2.4	665	2	US-08-883-795A-36	Sequence 36, Appl
20	51.2	2.4	1431	4	US-09-316-083-2	Sequence 2, Appl
21	51.2	2.4	3933	2	US-08-311-722-3	Sequence 1, Appl
22	51.2	2.4	3933	2	US-08-831-722-3	Sequence 1, Appl
23	51	2.4	615	4	US-08-416-186	Sequence 3, Appl
24	51	2.4	2539	3	US-08-849-522-3	Sequence 186, App
25	50.8	2.3	602	1	US-08-229-393-1	Sequence 1, Appl
26	50.6	2.3	8920	2	US-08-446-855A-1	Sequence 1, Appl
27	50.6	2.3	8920	4	US-09-150-741-1	Sequence 1, Appl

SUMMARIES

RESULTS

Result No.	Score	Query	Match Length	DB	ID	Description
1	78	3.6	7218	1	US-08-232-463-14	Sequence 14, Appl
2	61.4	2.8	837	4	US-08-998-416-288	Sequence 288, App
3	57.2	3.0	3095	6	S231168-1	Patent No. 567367
4	57	2.6	837	4	US-08-998-416-288	Sequence 288, App
5	56.4	2.6	615	4	US-08-998-16-1137	Sequence 186, App
6	56.2	2.6	636	4	US-08-998-16-1137	Sequence 186, App
7	56	2.6	19124	2	US-08-4-826B-13	Sequence 13, Appl
8	54.8	2.5	636	4	US-08-998-16-1137	Sequence 1137, Appl
9	54.2	2.5	1186	2	US-08-7-311-722-5	Sequence 5, Appl
10	53.8	2.5	6243	2	US-09-056-075-1	Sequence 1, Appl
11	53.6	2.5	19124	2	US-08-887-826B-13	Sequence 13, Appl
12	53.2	2.5	3138	1	US-07-867-106-4	Sequence 4, Appl
13	52.6	2.4	665	2	US-08-883-795A-36	Sequence 36, Appl
14	52.2	2.4	1826	4	US-09-686-601-11	Sequence 11, Appl
15	52.2	2.4	1826	4	US-09-687-47-11	Sequence 11, Appl
16	52	2.4	8920	2	US-08-446-855A-1	Sequence 1, Appl
17	52	2.4	8920	4	US-09-150-741-1	Sequence 1, Appl
18	51.8	2.4	782	4	US-09-007-119-15	Sequence 15, Appl
19	51.2	2.4	665	2	US-08-883-795A-36	Sequence 36, Appl
20	51.2	2.4	1431	4	US-09-316-083-2	Sequence 2, Appl
21	51.2	2.4	3933	2	US-08-311-722-3	Sequence 1, Appl
22	51.2	2.4	3933	2	US-08-831-722-3	Sequence 1, Appl
23	51	2.4	615	4	US-08-416-186	Sequence 3, Appl
24	51	2.4	2539	3	US-08-849-522-3	Sequence 186, App
25	50.8	2.3	602	1	US-08-229-393-1	Sequence 1, Appl
26	50.6	2.3	8920	2	US-08-446-855A-1	Sequence 1, Appl
27	50.6	2.3	8920	4	US-09-150-741-1	Sequence 1, Appl

ALIGNMENTS

RESULTS

Result No.	Score	Query	Match Length	DB	ID	Description
1	78	3.6	7218	1	US-08-232-463-14	Sequence 14, Appl
2	61.4	2.8	837	4	US-08-998-416-288	Sequence 288, App
3	57.2	3.0	3095	6	S231168-1	Patent No. 567367
4	57	2.6	837	4	US-08-998-416-288	Sequence 288, App
5	56.4	2.6	615	4	US-08-998-16-1137	Sequence 186, App
6	56.2	2.6	636	4	US-08-998-16-1137	Sequence 186, App
7	56	2.6	19124	2	US-08-4-826B-13	Sequence 13, Appl
8	54.8	2.5	636	4	US-08-998-16-1137	Sequence 1137, Appl
9	54.2	2.5	1186	2	US-08-7-311-722-5	Sequence 5, Appl
10	53.8	2.5	6243	2	US-09-056-075-1	Sequence 1, Appl
11	53.6	2.5	19124	2	US-08-887-826B-13	Sequence 13, Appl
12	53.2	2.5	3138	1	US-07-867-106-4	Sequence 4, Appl
13	52.6	2.4	665	2	US-08-883-795A-36	Sequence 36, Appl
14	52.2	2.4	1826	4	US-09-686-601-11	Sequence 11, Appl
15	52.2	2.4	1826	4	US-09-687-47-11	Sequence 11, Appl
16	52	2.4	8920	2	US-08-446-855A-1	Sequence 1, Appl
17	52	2.4	8920	4	US-09-150-741-1	Sequence 1, Appl
18	51.8	2.4	782	4	US-09-007-119-15	Sequence 15, Appl
19	51.2	2.4	665	2	US-08-883-795A-36	Sequence 36, Appl
20	51.2	2.4	1431	4	US-09-316-083-2	Sequence 2, Appl
21	51.2	2.4	3933	2	US-08-311-722-3	Sequence 1, Appl
22	51.2	2.4	3933	2	US-08-831-722-3	Sequence 1, Appl
23	51	2.4	615	4	US-08-416-186	Sequence 3, Appl
24	51	2.4	2539	3	US-08-849-522-3	Sequence 186, App
25	50.8	2.3	602	1	US-08-229-393-1	Sequence 1, Appl
26	50.6	2.3	8920	2	US-08-446-855A-1	Sequence 1, Appl
27	50.6	2.3	8920	4	US-09-150-741-1	Sequence 1, Appl

INFORMATION FOR SEQ ID NO: 14:

SEQUENCE CHARACTERISTICS:

LENGTH: 7218 base Pairs

TYPE: nucleic acid

STRANDEDNESS: single

TOPOLOGY: linear

IMMEDIATE SOURCE:

CLONE: pTZgpt-F1s

SEQUENCE 1: US-08-232-463-14

sequence 15, Appl
sequence 2, Appl
sequence 14, Appl
sequence 20, Appl
sequence 937, Appl
sequence 8, Appl
sequence 8, Appl
sequence 8, Appl
sequence 1, Appl
sequence 1, Appl
sequence 1, Appl
sequence 4, Appl
sequence 1, Appl
patent No. 5,213,974
Sequence 1, Appl
Sequence 14, Appl
Sequence 32, Appl
Sequence 4, Appl

STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: cDNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
US-08-487-826B-13

REGISTRATION NUMBER: 38,241
REFERENCE/DOCKET NUMBER: PFF/
TELECOMMUNICATION INFORMATION
TELEPHONE: 919-441-8857
TELEFAX: 919-541-8689
INFORMATION FOR SEQ ID NO: 1137
SEQUENCE CHARACTERISTICS:
LENGTH: 636 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
ORIGINAL SOURCE:
ORGANISM: PAG1692RP
US-08-998-416-1137

RESULT 8 US-08-998-416-1137
; Sequence 11/7, Application US/08998416
; Patent No. 623924

GENERAL INFORMATION:

APPLICANT: Philippson, Peter
APPLICANT: Pohlmann, Rainer
APPLICANT: Steiner, Sabine
APPLICANT: Mohr, Christine
APPLICANT: Wendland, Jürgen
APPLICANT: Knechtli, Philipp
APPLICANT: Rabitschung, Corinne

TITLE OF INVENTION: GENOMIC DNA SEQUENCES OF ASHBYA GOSSYPII
TITLE OF INVENTION: AND USES THEREOF

NUMBER OF SEQUENCES: 1152

CORRESPONDENCE ADDRESS:

ADDRESSEE: No. 623924 Artis Corporation
STREET: 3054 Cornwallis Road
CITY: Research Triangle Park
STATE: No. 6239264th Carolina
COUNTRY: USA
ZIP: 27709

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/998,416
FILING DATE: 24-DEC-1997
CLASSIFICATION: 435

PRIOR APPLICATION DATA:

APPLICATION NUMBER: CH 0016/97
FILING DATE: 31-DEC-1996

ATTORNEY/AGENT INFORMATION:

NAME: Meigs, J. Timothy

RESULT 9 US-08-731-722-5/C
; Sequence 5, Application US/08731722
; Patent No. 5951971

GENERAL INFORMATION:

APPLICANT: Martin, Frank N.

TITLE OF INVENTION: Biocontrol of Fungal Soilborne Pathogens

TITLE OF INVENTION: by Pythium oligandrum

NUMBER OF SEQUENCES: 8

CORRESPONDENCE ADDRESS:

ADDRESSEE: Saliwanchik & Saliwanchik
STREET: 2421 N.W. 41st Street, Suite A-1
CITY: Gainesville
STATE: FL
COUNTRY: US
ZIP: 32605-6669

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/731,722
FILING DATE: 24-DEC-1997
CLASSIFICATION: 424

ATTORNEY/AGENT INFORMATION:

NAME: Whillock, Ted W.

Search completed: January 24, 2002, 22:05:56
Job time: 1090 sec